



FF6

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/069,635A

Source:

PCT

Date Processed by STIC:

2/11/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT

MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/069,635A

DATE: 02/11/2004

TIME: 11:21:41

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\02112004\J069635A.raw

3 <110> APPLICANT: The Procter & Gamble Company Dykstra, Robert Miracle, Gregory
 5 <120> TITLE OF INVENTION: Controlled Availability of Formulation Components,
 Compositions and
 6 Laundry Methods Employing Same
 8 <130> FILE REFERENCE: 7749X
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/069,635A
 C--> 10 <141> CURRENT FILING DATE: 2002-09-20
 10 <150> PRIOR APPLICATION NUMBER: 60/151,002
 W--> 11 <150> PRIOR APPLICATION NUMBER: 1999-08-27<150> 60/151,004<151> 1999-08-27
 13 <160> NUMBER OF SEQ ID NOS: 18
 15 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 21
 19 <212> TYPE: DNA
 20 <213> ORGANISM: Aspergillus aculeatus
 22 <400> SEQUENCE: 1

E--> 23 attcatttgt ggacagtgga c

24 21

27 <210> SEQ ID NO: 2

28 <211> LENGTH: 20

29 <212> TYPE: DNA

30 <213> ORGANISM: Aspergillus aculeatus

32 <400> SEQUENCE: 2

E--> 33 gttgatcgca cattgaacca

34 20

37 <210> SEQ ID NO: 3

38 <211> LENGTH: 20

39 <212> TYPE: DNA

40 <213> ORGANISM: Aspergillus aculeatus

42 <400> SEQUENCE: 3

E--> 43 accccagccg accgattgtc

44 20

47 <210> SEQ ID NO: 4

48 <211> LENGTH: 20

49 <212> TYPE: DNA

50 <213> ORGANISM: Aspergillus aculeatus

52 <400> SEQUENCE: 4

E--> 53 cttccttacc tcaccatcat

54 20

57 <210> SEQ ID NO: 5

Does Not Comply
 Corrected Diskette Needed (pg 15)

21 - please insert

20 - please insert

Same error

Same error

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TIME: 11:21:41

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\02112004\J069635A.raw

58 <211> LENGTH: 20
59 <212> TYPE: DNA
60 <213> ORGANISM: Aspergillus aculeatus
62 <400> SEQUENCE: 5
E--> 63 ttaacatctt ttcacatga
64 20
67 <210> SEQ ID NO: 6
68 <211> LENGTH: 20
69 <212> TYPE: DNA
70 <213> ORGANISM: Aspergillus aculeatus
72 <400> SEQUENCE: 6
E--> 73 agctttccct tctctccctt
74 20
77 <210> SEQ ID NO: 7
78 <211> LENGTH: 28
79 <212> TYPE: DNA
80 <213> ORGANISM: Aspergillus aculeatus
82 <400> SEQUENCE: 7
E--> 83 gccaccctgg cttccgctgc cagcctcc
84 28
87 <210> SEQ ID NO: 8
88 <211> LENGTH: 20
89 <212> TYPE: DNA
90 <213> ORGANISM: Aspergillus aculeatus
92 <400> SEQUENCE: 8
E--> 93 gacagtagca atccagcatt
94 20
97 <210> SEQ ID NO: 9
98 <211> LENGTH: 20
99 <212> TYPE: DNA
100 <213> ORGANISM: Aspergillus aculeatus
102 <400> SEQUENCE: 9
E--> 103 agcatcagcc gctttgtaca
104 20
107 <210> SEQ ID NO: 10
108 <211> LENGTH: 20
109 <212> TYPE: DNA
110 <213> ORGANISM: Aspergillus aculeatus
112 <400> SEQUENCE: 10
E--> 113 ccatgaagtt caccgtattg
114 20
117 <210> SEQ ID NO: 11
118 <211> LENGTH: 20
119 <212> TYPE: DNA
120 <213> ORGANISM: Aspergillus aculeatus
122 <400> SEQUENCE: 11
E--> 123 gcactgcttc tctcccaggt
124 20
127 <210> SEQ ID NO: 12

same errors

RAW SEQUENCE LISTING

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TIME: 11:21:41

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\02112004\J069635A.raw

128 <211> LENGTH: 20
 129 <212> TYPE: DNA
 130 <213> ORGANISM: Aspergillus aculeatus
 132 <400> SEQUENCE: 12
 E--> 133 gtgggcggcc cctcaggcaa
 134 20
 137 <210> SEQ ID NO: 13
 138 <211> LENGTH: 20
 139 <212> TYPE: DNA
 140 <213> ORGANISM: Aspergillus aculeatus
 142 <400> SEQUENCE: 13
 E--> 143 acgtcctcc aattttctct
 144 20
 147 <210> SEQ ID NO: 14
 148 <211> LENGTH: 19
 149 <212> TYPE: DNA
 150 <213> ORGANISM: Aspergillus aculeatus
 152 <400> SEQUENCE: 14
 E--> 153 ggctggtagt aatgagtct
 154 19
 157 <210> SEQ ID NO: 15
 158 <211> LENGTH: 20
 159 <212> TYPE: DNA
 160 <213> ORGANISM: Aspergillus aculeatus
 162 <400> SEQUENCE: 15
 E--> 163 ggcgcagagt ttggccaggc
 164 20
 167 <210> SEQ ID NO: 16
 168 <211> LENGTH: 21
 169 <212> TYPE: DNA
 170 <213> ORGANISM: Aspergillus aculeatus
 172 <400> SEQUENCE: 16
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 174 21
 177 <210> SEQ ID NO: 17
 178 <211> LENGTH: 347
 179 <212> TYPE: DNA
 180 <213> ORGANISM: Aspergillus aculeatus
 182 <400> SEQUENCE: 17
 E--> 183 aaagattcat ttgtggacag tggacgttg tgcacattg aaccaacccc agccgaccga 60
 184 60 move to
 E--> 186 ttgtccttcc ttacctcacc atcatttaac atcttttcac catgaagctt tcccttctct 120
 187 120 move to
 E--> 189 cccttgccac cctggcttcc gctgccagcc tccagcgcgc cacacttctg cggtcagtgg 180
 190 180 move to
 E--> 192 gataccgcca ccgccggtga cttcaccctg tacaacgacc tttggggcga gacggccggc 240
 193 240 move to
 E--> 195 accggctccc agtgcactgg agtcgactcc tacagcggcg acaccatcgc ttgtcacacc 300
 196 300 move to

Same errors

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TIME: 11:21:41

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\02112004\J069635A.raw

E--> 198 agcaggtcct ggtcggagta gcagcagcgt caagagctat gccaacg 347
199 347
202 <210> SEQ ID NO: 18
203 <211> LENGTH: 294
204 <212> TYPE: DNA
205 <213> ORGANISM: Aspergillus aculeatus
207 <400> SEQUENCE: 18
E--> 208 cagcatctcc attgagtaat cacgttggtg ttcggtggcc cgccgtggtg cgtggcggag
209 60
E--> 211 gctgccggga gacgggtggg gatggtggtg ggagagaatg tagggcgccg tgtttcagtc
212 120
E--> 214 cctaggcagg ataccgaaa accgtgtggt aggaggttta taggtttcca ggagacgctg
215 180
E--> 217 tataggggat aaatgagatt gaatggtggc cacactcaaa ccaaccaggt cctgtacata
218 240
E--> 220 caatgcatat accaattata cctacaaaaa aaaaaaaaaa aaaaaaaaaa aaaa
221 294

Some
errors

VERIFICATION SUMMARY

DATE: 02/11/2004

PATENT APPLICATION: US/10/069,635A

TIME: 11:21:42

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\02112004\J069635A.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:11 M:289 W: Identifier Missing or Out-Of-Order, <150> PRIOR APP FILING DATE
L:23 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:21 SEQ:1
L:33 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:2
L:43 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:3
L:53 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:4
L:63 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:5
L:73 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:6
L:83 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:28 SEQ:7
L:93 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:8
L:103 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:9
L:113 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:10
L:123 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:11
L:133 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:12
L:143 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:13
L:153 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:19 SEQ:14
L:163 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:20 SEQ:15
L:173 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:21 SEQ:16
L:183 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:17
M:254 Repeated in SeqNo=17
L:208 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:60 SEQ:18
M:254 Repeated in SeqNo=18